

*In the claims:*

Please amend claims 23, 25, 27, 29, 33-36, 39, 40, 43, 44, 47, 48, 60-67, 72-74, 77, 78, 81, 82, 85 and 86 as follows.

D<sup>2</sup> ~~3.~~<sub>23.</sub> (Once Amended) The polypeptide of claim ~~22.~~<sub>2.</sub> wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>3</sup> ~~5.~~<sub>25.</sub> (Once Amended) The polypeptide of claim ~~24.~~<sub>4.</sub> wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>4</sup> ~~7.~~<sub>27.</sub> (Once Amended) The polypeptide of claim ~~26.~~<sub>6.</sub> wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>5</sup> ~~9.~~<sub>29.</sub> (Once Amended) The polypeptide of claim ~~28.~~<sub>8.</sub> wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>6</sup> ~~10.~~<sub>30.</sub> (Once Amended) The polypeptide of claim ~~22.~~<sub>2.</sub> wherein the amino acid sequence is fused to a heterologous polypeptide.

D<sup>7</sup> ~~11.~~<sub>31.</sub> (Twice Amended) An isolated polypeptide comprising a first amino acid sequence that is at least 95% identical to a second amino acid sequence selected from the group consisting of:

(a) amino acids 1 to 381 of SEQ ID NO:2;

(b) amino acids 2 to 381 of SEQ ID NO:2;

(c) amino acids 25 to 381 of SEQ ID NO:2; and

(d) a polypeptide fragment of SEQ ID NO:2,

wherein said polypeptide or polypeptide fragment stimulates cellular proliferation.

*D<sup>7</sup> cont'd*  
~~12.~~ ~~35.~~ (Once Amended) The polypeptide of claim ~~34~~<sup>11</sup>, wherein said second amino acid sequence is (a).

*D<sup>8</sup>*  
~~13.~~ ~~36.~~ (Once Amended) The polypeptide of claim ~~35~~<sup>12</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>9</sup>*  
~~14.~~ ~~39.~~ (Once Amended) The polypeptide of claim ~~34~~<sup>11</sup>, wherein said second amino acid sequence is (b).

~~15.~~ ~~40.~~ (Once Amended) The polypeptide of claim ~~39~~<sup>14</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

*D<sup>10</sup>*  
~~16.~~ ~~43.~~ (Once Amended) The polypeptide of claim ~~34~~<sup>11</sup>, wherein said second amino acid sequence is (c).

~~17.~~ ~~44.~~ (Once Amended) The polypeptide of claim ~~43~~<sup>16</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

<sup>11</sup>  
~~18.~~ ~~47.~~ (Once Amended) The polypeptide of claim ~~34~~, wherein said second amino acid sequence is (d).

<sup>11</sup>  
~~19.~~ ~~48.~~ (Once Amended) The polypeptide of claim ~~47~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

~~21.~~ ~~60.~~ (Once Amended) The polypeptide of claim ~~59~~, wherein said amino acid sequence is (a).

~~22.~~ ~~61.~~ (Once Amended) The polypeptide of claim ~~60~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

<sup>12</sup>  
~~23.~~ ~~62.~~ (Once Amended) The polypeptide of claim ~~59~~, wherein said amino acid sequence is (b).

~~24.~~ ~~63.~~ (Once Amended) The polypeptide of claim ~~62~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

~~25.~~ ~~64.~~ (Once Amended) The polypeptide of claim ~~59~~, wherein said amino acid sequence is (c).

~~26.~~ ~~65.~~ (Once Amended) The polypeptide of claim ~~64~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

27.  
~~66.~~

20,  
~~59.~~

(Once Amended) The polypeptide of claim ~~59~~, wherein said amino acid sequence is (d).

28.  
~~67.~~

27  
~~68.~~

(Once Amended) The polypeptide of claim ~~68~~, wherein the amino acid sequence is fused to a heterologous polypeptide.

29.  
~~72.~~

(Twice Amended) An isolated polypeptide comprising a first amino acid sequence that is at least 95% identical to a second amino acid sequence selected from the group consisting of:

(a) the amino acid sequence of the full-length polypeptide encoded by the human cDNA contained in ATCC Deposit Number 75904;

(b) the amino acid sequence of the full-length polypeptide, lacking the N-terminal methionine, encoded by the human cDNA contained in ATCC Deposit Number 75904;

(c) the amino acid sequence of the mature polypeptide encoded by the human cDNA contained in ATCC Deposit Number 75904; and

(d) a polypeptide fragment of the polypeptide encoded by the human cDNA contained in ATCC Deposit Number 75904;

wherein said polypeptide or polypeptide fragment stimulates cellular proliferation.

30.  
~~73.~~

29,  
~~72.~~

(Once Amended) The polypeptide of claim ~~72~~, wherein said second amino acid sequence is (a).

D<sup>14</sup>

31.  
~~74.~~

(Once Amended) The polypeptide of claim ~~73~~<sup>30</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

32.  
~~77.~~

(Once Amended) The polypeptide of claim ~~72~~<sup>29</sup>, wherein said second amino acid sequence is (b).

33.  
~~78.~~

(Once Amended) The polypeptide of claim ~~77~~<sup>32</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

34.  
~~81.~~

(Once Amended) The polypeptide of claim ~~72~~<sup>29</sup>, wherein said second amino acid sequence is (c).

35.  
~~82.~~

(Once Amended) The polypeptide of claim ~~81~~<sup>34</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.

36.  
~~85.~~

(Once Amended) The polypeptide of claim ~~72~~<sup>29</sup>, wherein said second amino acid sequence is (d).

37.  
~~86.~~

(Once Amended) The polypeptide of claim ~~85~~<sup>36</sup>, wherein the amino acid sequence is fused to a heterologous polypeptide.